

SEQUENCE LISTING

<110> Sera, Takashi
 <120> Nuclear-Envelope and Nuclear-Lamina Binding Chimeras for
 Modulating Gene Expression
 <130> 109845-163
 <160> 18
 <170> PatentIn version 3.2
 <210> 1
 <211> 25
 <212> PRT
 <213> Artificial
 <220>
 <223> Zinc finger domain
 <220>
 <221> MISC_FEATURE
 <222> (2)..(5)
 <223> Amino acids 2-5 are Xaa wherein Xaa = any amino acid, and up to
 two amino acids can be missing.
 <220>
 <221> MISC_FEATURE
 <222> (7)..(18)
 <223> Xaa can be any amino acid
 <220>
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 <222> (20)..(24)
 <223> Amino acids 20-24 are Xaa wherein Xaa = any amino acid, and up to
 two amino acids can be missing.
 <400> 1
 Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Xaa His Xaa Xaa Xaa Xaa His
 20 25
 <210> 2
 <211> 32
 <212> PRT
 <213> Artificial
 <220>
 <223> Second zinc finger domain
 <220>
 <221> MISC_FEATURE
 <222> (1)..(3)
 <223> Xaa can be any amino acid
 <220>
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 <222> (5)..(8)
 <223> Amino acids 5-8 are Xaa wherein Xaa = any amino acid, and up to
 two amino acids can be missing

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<220>
<221> MISC_FEATURE
<222> (10)..(14)
<223> Amino acid 15 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr,
Met, Leu, Ile, Glu or Asp.

<220>
<221> misc_feature
<222> (15)..(15)
<223> Xaa can be any naturally occurring amino acid

<220>
<221> MISC_FEATURE
<222> (16)..(16)
<223> Xaa can be any amino acid

<220>
<221> MISC_FEATURE
<222> (17)..(17)
<223> Amino acid 17 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val,
Ala, Asp or Glu.

<220>
<221> MISC_FEATURE
<222> (18)..(18)
<223> Amino acid 18 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala,
Val, Thr, Asp, or Glu

<220>
<221> MISC_FEATURE
<222> (19)..(20)
<223> Xaa can be any amino acid

<220>
<221> MISC_FEATURE
<222> (21)..(21)
<223> Amino acid 21 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr,
Leu, Ile, Met, Glu or Asp.

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<221> MISC_FEATURE
<222> (23)..(27)
<223> Amino acids 23-27 are Xaa wherein Xaa = any amino acid, adn up to
two amino acids can be missing.

<220>
<221> MISC_FEATURE
<222> (29)..(32)
<223> Xaa can be any amino acid

<400> 2

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa
20 25 30

<210> 3
<211> 28
<212> PRT
<213> Artificial

<220>
<223> Zinc finger domain

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<220>
 <221> MISC_FEATURE
 <222> (13)..(13)
 <223> Amino acid 13 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr, Met, Leu, Ile, Glu or Asp.

<220>
 <221> MISC_FEATURE
 <222> (15)..(15)
 <223> Amino acid 15 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val, Ala, Asp or Glu.

<220>
 <221> MISC_FEATURE
 <222> (16)..(16)
 <223> Amino acid 16 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala, Val, Thr, Asp or Glu.

<220>
 <221> MISC_FEATURE
 <222> (19)..(19)
 <223> Amino acid 19 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr, Leu, Ile, Met, Glu or Asp.

<400> 3

Pro Tyr Lys Cys Pro Glu Cys Gly Lys Ser Phe Ser Xaa Ser Xaa Xaa
 1 5 10 15

Leu Gln Xaa His Gln Arg Thr His Thr Gly Glu Lys
 20 25

<210> 4
 <211> 5
 <212> PRT
 <213> Artificial

<220>
 <223> peptide

<400> 4

Gly Gly Gly Gly Ser
 1 5

<210> 5
 <211> 11
 <212> PRT
 <213> Human immunodeficiency virus

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 <221> MISC_FEATURE
 <222> (1)..(11)
 <223> HIV Tat protein domain

<400> 5

Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
 1 5 10

<210> 6
 <211> 9
 <212> DNA
 <213> Human immunodeficiency virus

<220>
 <221> misc_feature
 <222> (1)..(9)
 <223> HIV DNA Binding Domain

<400> 6
 gcagaagcc

9

<210> 7
 <211> 19
 <212> DNA
 <213> Artificial

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 <223> DNA target sequence

<400> 7
 gtgtgggtga gtgagtgtg

19

<210> 8
 <211> 19
 <212> DNA
 <213> Artificial

<220>
 <223> DNA target sequence

<400> 8
 ggggctgggg gcggtgtct

19

<210> 9
 <211> 7
 <212> PRT
 <213> Simian virus 40

<220>
 <221> MISC_FEATURE
 <222> (1)..(7)
 <223> Peptide from SV40 large T antigen

<400> 9

Pro Lys Lys Lys Arg Lys Val
 1 5

<210> 10
 <211> 16
 <212> PRT
 <213> Artificial

<220>
 <223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

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 <221> MISC_FEATURE
 <222> (1)..(16)

<223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

<400> 10

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
1 5 10 15

<210> 11

<211> 34

<212> PRT

<213> Herpes Simplex Virus

<220>

<221> MISC_FEATURE

<222> (1)..(34)

<223> Residues 267-300 of the HSV VP22 protein

<400> 11

Asp Ala Ala Thr Ala Thr Arg Gly Arg Ser Ala Ala Ser Arg Pro Thr
1 5 10 15

Glu Arg Pro Arg Ala Pro Ala Arg Ser Ala Ser Arg Pro Arg Arg Pro
20 25 30

Val Glu

<210> 12

<211> 11

<212> PRT

<213> Artificial

<220>

<223> Basic peptide with cellular uptake activity

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<221> MISC_FEATURE

<222> (1)..(11)

<223> Basic peptide with cellular uptake activity

<400> 12

Tyr Ala Arg Ala Ala Ala Arg Gln Ala Arg Ala
1 5 10

<210> 13

<211> 9

<212> PRT

<213> Artificial

<220>

<223> Basic peptide with cellular uptake activity, "R9"

<400> 13

Arg Arg Arg Arg Arg Arg Arg Arg
1 5

<210> 14
 <211> 16
 <212> PRT
 <213> Artificial

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 <223> D-penetratin peptide

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 <221> MISC_FEATURE
 <222> (1)..(16)
 <223> All amino acids are in the D-form

 <400> 14
 Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
 1 5 10 15

 <210> 15
 <211> 16
 <212> PRT
 <213> Artificial

 <220>
 <223> Peptide Syn B1 from Antennapedia homeodomain protein

 <400> 15
 Arg Gly Gly Arg Leu Ser Tyr Ser Arg Arg Arg Phe Ser Thr Ser Thr
 1 5 10 15

 <210> 16
 <211> 10
 <212> PRT
 <213> Artificial

 <220>
 <223> L-SynB3 peptide from Antennapedia homeodomain protein

 <400> 16
 Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe
 1 5 10

 <210> 17
 <211> 10
 <212> PRT
 <213> Artificial

 <220>
 <223> D-SynB3 peptide from Antennapedia homeodomain pro

 <220>
 <221> MISC_FEATURE
 <222> (1)..(10)
 <223> All amino acids are in the D-form

 <400> 17
 Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe
 1 5 10

<210> 18

<211> 8

<212> PRT

<213> Artificial

<220>

<223> Flag Epitope Peptide

<400> 18

Asp Tyr Lys Asp Asp Asp Asp Lys
1 5